SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Frank, Bruce L. Roberts, Peter C. Goodchild, John Craig, J. Charles Mills, John S. Slade, Andrew Roberts, Noel A. Jupp, Raymond
- (ii) TITLE OF INVENTION: OLIGONUCLEOTIDES SPECIFIC FOR HEPATITIS B VIRUS
- (iii) NUMBER OF SEQUENCES: 56
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Lappin & Kusmer
 - (B) STREET: 200 State Street
 - (C) CITY: Boston
 - (D) STATE: MA
 - (E) COUNTRY: USA (F) ZIP: 02109
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk

 - (B) COMPUTER: IBM PC compatible (C) OPERATING SYSTEM: PC-DOS/MS-DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
- (viii) ATTORNEY/AGENT INFORMATION: (A) NAME: Kerner, Ann-Louise
 - (B) REGISTRATION NUMBER: 33,523
 - (C) REFERENCE/DOCKET NUMBER: HYZ-041
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 617-330-1300
 - (B) TELEFAX: 617-330-1311
- (2) INFORMATION FOR SEO ID NO:1:
 - (i) SEQUENCE CHARACTERISTICS:

	(A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:1:	
GGTGCGCA	GA CCAATTTATG	20
(2) INFO	RMATION FOR SEQ ID NO:2:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 19 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:2:	
CATGGTGC'	TG GTGCGCAGA	19
(2) INFO	RMATION FOR SEQ ID NO:3:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:3:	
GAAAAAGT"	TG CATGGTGCTG	20

(2)	INFO	RMATION FOR SEQ ID NO:4:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:4:	
GAG	GTGAA	AA AGTTGCATGG	20
(2)	INFO	RMATION FOR SEQ ID NO:5:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:5:	
AGG	CAGAG	GT GAAAAAGTTG	20
(2)	INFO	RMATION FOR SEQ ID NO:6:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
	(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:6:	

AGGCAGAGGT GA	12
(2) INFORMATION FOR SEQ ID NO:7:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	1
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: YES	
(xi) SEQUENCE DESCRIPTION: SEQ	ID NO:7:
AGAGATGATT AGGCAGAGGT	20
(2) INFORMATION FOR SEQ ID NO:8:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: YES	
(xi) SEQUENCE DESCRIPTION: SEQ	ID NO:8:
GACATGAACA AGAGATGATT AGGCAGAGGT	30
(2) INFORMATION FOR SEQ ID NO:9:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	3
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: YES	

(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:9:	
GACATGTAC	CA AGAGATGATT	20
(2) INFOR	RMATION FOR SEQ ID NO:10:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:10:	
GTAGGACAT	TG AACAAGAGAT	20
(2) INFOR	RMATION FOR SEQ ID NO:11:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:11:	
TTGGAGGCT	T GAACAGTAGG	20
(2) INFOR	RMATION FOR SEQ ID NO:12:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	

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(iv) ANTI-SENSE: YES	•
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
CACAGCTTGG AGGCTTGAAC	20
(2) INFORMATION FOR SEQ ID NO:13:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: YES	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:	
AGCCACCCAA GGCACAGCTT	20
(2) INFORMATION FOR SEQ ID NO:14:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDBESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: YES	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:	
TCGATGTCCA TGCCCCAAAG	20
(2) INFORMATION FOR SEQ ID NO:15:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	

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(iii	HYPOTHETICAL: NO	
(iv	ANTI-SENSE: YES	
(xi	SEQUENCE DESCRIPTION: SEQ ID NO:15:	
TAAGGGT	CGA TGTCCATGCC	20
(2) INF	ORMATION FOR SEQ ID NO:16:	
(i	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii	MOLECULE TYPE: DNA	
(iii	HYPOTHETICAL: NO	
(iv	ANTI-SENSE: YES	
(xi	SEQUENCE DESCRIPTION: SEQ ID NO:16:	
TTATAAG	GGT CGATGTCCAT	20
(2) INF	ORMATION FOR SEQ ID NO:17:	
(i	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii	MOLECULE TYPE: DNA	
(iii	HYPOTHETICAL: NO	
(iv	ANTI-SENSE: YES	
(xi	SEQUENCE DESCRIPTION: SEQ ID NO:17:	
AAATTCT	TTA TAAGGGTCGA TGTCCAT	27
(2) INF	DRMATION FOR SEQ ID NO:18:	
(i	SEQUENCE CHARACTERISTICS: (A) LENGTH: 12 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

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(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:18:	
TATAAGGG'	TC GA	12
(2) INFO	RMATION FOR SEQ ID NO:19:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:19:	
AAATTCTT'	TA TAAGGGTCGA	20
(2) INFO	RMATION FOR SEQ ID NO:20:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:20:	
GTATCTAG	AA GATCTCGTAC	20
(2) INFO	RMATION FOR SEQ ID NO:21:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid	

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	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:21:	
GCGGTGTC	TA GAAGATCTCG	20
(2) INFO	RMATION FOR SEQ ID NO:22:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:22:	
GAGGCGGT	GT CTAGGAGATC	20
(2) INFO	RMATION FOR SEQ ID NO:23:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDENDESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:23:	
GAGCTGAG	GC GGTGTCTAGG	20
(2) INFO	RMATION FOR SEQ ID NO:24:	

(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:24:	
ATACAGAG	CT GAGGCGGTAT	20
(2) INFO	RMATION FOR SEQ ID NO:25:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:25:	
TCCCGATA	CA GAGCTGAGGC	20
(2) INFO	RMATION FOR SEQ ID NO:26:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:26:	
AGGCTTCC	CG ATACAGAGCT	20

(2) INFO	RMATION FOR SEQ ID NO:27:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:27:	
ACAATGCT	CA GGAGACTCTA	20
(2) INFO	RMATION FOR SEQ ID NO:28:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:28:	
GCAGTATG	GT GAGGTGAGCA	20
(2) INFO	RMATION FOR SEQ ID NO:29:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:29:	

GAGTGCAGTA TGGTGAGGTG	20
(2) INFORMATION FOR SEQ ID NO:30:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: YES	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:	
TGCCTGAGTG CAGTATGGTG	20
(2) INFORMATION FOR SEQ ID NO:31:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: YES	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:	
TTGCTTGCCT GAGTGCAGTA	20
(2) INFORMATION FOR SEQ ID NO:32:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE, VES	

(x:	i) SEQUENCE DESCRIPTION: SEQ ID NO:32:	
TAAGGG	TCGA AGAGATGATT	20
(2) IN	FORMATION FOR SEQ ID NO:33:	
(:	i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(i:	i) MOLECULE TYPE: DNA	
(ii	i) HYPOTHETICAL: NO	
(i	v) ANTI-SENSE: YES	
(x:	i) SEQUENCE DESCRIPTION: SEQ ID NO:33:	
AGAGATO	GATT TAAGGGTCGA	20
(2) IN	FORMATION FOR SEQ ID NO:34:	
(:	i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(i:	i) MOLECULE TYPE: DNA	
(ii	i) HYPOTHETICAL: NO	
(i	v) ANTI-SENSE: YES	
(x:	i) SEQUENCE DESCRIPTION: SEQ ID NO:34:	
TAAGGG	TCGA AGGCAGAGGT	20
(2) IN	FORMATION FOR SEQ ID NO:35:	
(.	i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(i	i) MOLECULE TYPE: DNA	
(ii	i) HYPOTHETICAL: NO	

(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:35:	
AGGCAGAGG	GT TAAGGGTCGA	20
(2) INFOR	RMATION FOR SEQ ID NO:36:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:36:	
TATAAGGGT	TC GAAGGCAGAG GTGA	24
(2) INFOR	RMATION FOR SEQ ID NO:37:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:37:	
AGGCAGAGG	GT GATATAAGGG TCGA	24
(2) INFOR	RMATION FOR SEQ ID NO:38:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	

(iii) HYPOTHETICAL: NO (iv) ANTI-SENSE: YES (xi) SEQUENCE DESCRIPTION: SEQ ID NO:38: AGAGATGATT AGGCAGAGGT TAAGGGTCGA (2) INFORMATION FOR SEQ ID NO:39: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	30
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:38: AGAGATGATT AGGCAGAGGT TAAGGGTCGA (2) INFORMATION FOR SEQ ID NO:39: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	30
AGAGATGATT AGGCAGAGGT TAAGGGTCGA (2) INFORMATION FOR SEQ ID NO:39: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	30
(2) INFORMATION FOR SEQ ID NO:39: (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	30
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	
(A) LENGTH: 30 base pairs(B) TYPE: nucleic acid(C) STRANDEDNESS: single	

(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: YES	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:	
GACATGAACA AGAGATGATT TAAGGGTCGA	30
(2) INFORMATION FOR SEQ ID NO:40:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: YES	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:	
AGAGATGATT TAAGGGTCGA TGTCCATGCC	30
(2) INFORMATION FOR SEQ ID NO:41:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 30 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	

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(ii)	MOLECULE TYPE: DNA	
(i	ii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(:	xi)	SEQUENCE DESCRIPTION: SEQ ID NO:41:	
AGGCA	GAGG	T TAAGGGTCGA TGTCCATGCC	30
(2) I	NFOF	MATION FOR SEQ ID NO:42:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(i	ii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(:	xi)	SEQUENCE DESCRIPTION: SEQ ID NO:42:	
GGCAT	TTGG	T GGTCTATAAG	20
(2) I	NFOF	MATION FOR SEQ ID NO:43:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(i:	ii)	HYPOTHETICAL: NO	
(:	iv)	ANTI-SENSE: YES	
(:	xi)	SEQUENCE DESCRIPTION: SEQ ID NO:43:	
GATAG	GGGC	TA TTTGGTGGTC	20
(2) I	NFOR	MATION FOR SEQ ID NO:44:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid	•

	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:44:	
TGTTGATA	GG ATAGGGGCAT	20
(2) INFO	RMATION FOR SEQ ID NO:45:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:45:	
ACCCAAGG	CA CAGCTTGGAG	20
(2) INFO	RMATION FOR SEQ ID NO:46:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:46:	
GACAGGGG	CA TTTGGTGGTC	20
(2) INFO	RMATION FOR SEQ ID NO:47:	

(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:47:	
GATAGGGG	CC TTTGGTGGTC	20
(2) INFO	RMATION FOR SEQ ID NO:48:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:48:	
GATAGGGG	CA TTTGGTGCTC	20
(2) INFO	RMATION FOR SEQ ID NO:49:	
(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
(iv)	ANTI-SENSE: YES	
(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:49:	
GACAGGGG	CC TTTGGTGCTC	20

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(2)	INFO	RMATION FOR SEQ ID NO:50:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 20 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
((iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:50:	
ATGI	TTTT	GG CGTCTTCCAT	20
(2)	INFO	RMATION FOR SEQ ID NO:51:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 21 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
((iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: NO	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:51:	
TGCI	GGGG	GG AACTAATGAC T	21
(2)	INFO	RMATION FOR SEQ ID NO:52:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 25 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: NO	
	(xi)	SEQUENCE DESCRIPTION: SEO ID NO:52:	

GGAG	TGCG.	AA TCCACACTCC GAAAG	25
(2)	INFO	RMATION FOR SEQ ID NO:53:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
((iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:53:	
raag	GGTC	GA .	10
(2)	INFO	RMATION FOR SEQ ID NO:54:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 10 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: YES	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:54:	
AGGC	AGAG	FT	10
(2)	INFO	RMATION FOR SEQ ID NO:55:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 36 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: DNA	
(iii)	HYPOTHETICAL: NO	
	(iv)	ANTI-SENSE: NO	

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:55:	
ATATTCGCAC GAATTCAGCT TATAGACCAC CAAATG	
(2) INFORMATION FOR SEQ ID NO:56:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 34 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: DNA	
(iii) HYPOTHETICAL: NO	
(iv) ANTI-SENSE: NO	
(xi) SEOUENCE DESCRIPTION: SEO ID NO:56:	

GTTGGGATTG CTGCAGCTAT CTGGATTTGC GGTG

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36